

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 148592

TO: Vanessa L Ford Location: 3b25 / 3c18 Monday, March 28, 2005

Art Unit: 1645

Phone: 571-272-0857

Serial Number: 10 / 017168

From: Jan Delaval

Location: Biotech-Chem Library

Remsen 1a51

Phone: 571-272-22504

jan.delaval@uspto.gov

Search Notes	*		.) * i *-			
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From:

Chan, Christina

Sent: To:

Wednesday, March 23, 2005 8:02 AM Ford, Vanessa; STIC-Biotech/ChemLib RE: In re: 10/017,168 Sequence search

Subject:

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

-----Original Message-----

From:

Ford, Vanessa

Sent:

Tuesday, March 22, 2005 7:04 PM

To:

Chan, Christina

Subject:

In re: 10/017,168 Sequence search

Please search SEQ ID NOs: 1, 3,5,19, 21, 23 and 25. Please include interference searches. Please rush.

Vanessa L. Ford

Biotechnology Patent Examiner Office: REM 3B25

Mailbox: REM 3C18 Phone: 571.272.0857

Art unit:1645

STAFF USE ONLY Searcher: Searcher Phone: 2- でいく Date Searcher Picked up: 3123 Date Completed: 3/2016 Searcher Prep/Rev. Time: 20 Online Time:__

Type of Search

NA#:__C Interference: Oligomer:_ S/L:_ Encode/Transl: Structure#:

Inventor:___

Litigation:_

Vendors and cost where applicable STN:_ DIALOG: QUESTEL/ORBIT: LEXIS/NEXIS:_ SEQUENCE SYSTEM: WWW/Internet:_ Other(Specify):_

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RESULT 1
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Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological sample.
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                  Claim 19; Fig 5; 73pp; English
                                                                                       WPI; 2001-080711/09.
P-PSDB; AAB48316.
                                                                                                                    Liu H,
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                                                                                                                                                                                                                                                                                 Treponema pallidum; ssp. pallidum.
                                                                                                                                                                                                                                                                                                    Treponema pallidum; acidic repeat protein; arp; immunogenic; syphilis; yaws; bejel; ds.
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20-APR-2001
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The invention relates to a method of detecting presence of Treponema

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JOURNAL COMMENT REFERENCE AUTHORS SOURCE ORGANISM ACCESSION VERSION KEYWORDS RESULT 1 CL079800/c LOCUS DEFINITION ORIGIN FEATURES Query Match 6.9%; Best Local Similarity 51.4%; Matches 436; Conservative TITLE source Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Mesobatrachia; Pipcidea; Pipidae; Xenopodinae; Xenopus; Silurana. 1 (bases 1 to 1494) Kremitzki,C., Carter,J., McPherson,J., Warren,W., Graves,T., Mardis,E. and Wilson,R. A physical map of the xenopus tropicalis genome Unpublished (2003) Washington University School of Medicine Email: submissions@watson.wastl.edu Insert Length: 175000 Std Error: 0.00 Seq primer: Sp5 atctgccgtttcgatcct Class: BAC ends GSS. CL079800 1484 bp DNA linear GSS 31 CH216-156023_Sp5.1 CH216 Xenopus tropicalis genomic clone CH216-156023, genomic survey sequence. Genome Sequencing Center Contact: Richard K Wilson Xenopus tropicalis CL079800.1 GI:40535713 CL079800 Kenopus tropicalis (western clawed frog) quality sequence start: 549 quality sequence stop: 627. Location/Qualifiers /cell_line="Stock 248 F7A2, inbred N'" /clone_lib="CH216" /clone_e="Vector: pTARBAC2.1; CHORI-216 Xenopus tropicalis BAC library" /organism="Xenopus tropicalis" /mol_type="genomic DNA" /strain="Nigerian frog" /db xref="taxon:8364" /clone="CH216-156023" 1. .1484 sex="male" 0 Score 203.2; DB 9; Pred. No. 4.9e-44; 0; Mismatches 413; Indels Length 1484; GSS 31-DEC-2003 0 Gaps 0

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ORIGIN	CDS	REFERENCE AUTHORS TITLE JOURNAL FEATURES BOUICE	VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 AX068032 LOCUS DEFINITION
(Nichols strain) /(codon_start=1) /(codon_start=1) /(codon_start=1) /(trans1_table=1) /(trans1_table=1) /(trans1_table=1) /(trans1_table=1) /(db_xref="G1:12329841" /(db_xref="G1:12329841" /(db_xref="G1:12329841" /(trans1ation="MyvrSDMFPKNTAVEISNLEKNAKAQAVVIGHAGIPGLLVSLAP /(trans1ation="MyvrSDMFPKNTAVEISNLEKNAKAQAVVIGHAGIPGLLVSLAP /(trans1ation="MyvrSDMFPKNTAVEISNLEKNAKAQAVVIGHARPAQRDPLSSPP AAAAQLGIGVYQAVRVRVRTHGTVRCGSQTSQDLELSLASLPSRVPARPARPAQRDPLSSPP AAAAQLGIGVYQAVRVRVRTHGTVRCGSQTSQDJRKVVEPASEREGGEREVEDVPKVVEP AGRICGGEREVEDVPKVVEPASEREGGEREVEDAPKVV ASEREGGEREVENVPKVVEPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPEPASEREGGEREVEDVPKVVEPASEREGGEREVEDVPKVVEPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPASEREGGEREVEDVPKVPPASEREGGEREVE	/Organisms interviews particum /mol_type="unassigned DNA" /db_xref="taxon:160" 9192217 /note="unnamed protein product; Subspecies: pallidum	Liu,H., Steiner,B. and Rhodes,B. Liu,H., Steiner,B. and Rhodes,B. Compositions and methods for detecting Treponema pallidum Patent: WO 0077486-A 1 21-DEC-2000; THE GOVERNMENT OF THE UNITED STATES OF AMERICA (US) Location/Qualifiers 1. 2945 1. 2945	AX068032.1 GI:12329840 Treponema pallidum Treponema pallidum Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.	AX068032 2945 bp DNA linear PAT 21-JAN-2001 Sequence 1 from Patent WO0077486.

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Query Match 100.0%; Score 2945; Best Local Similarity 100.0%; Pred. No. 0; Matches 2945; Conservative 0; Mismatches

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Aac26603 Polymucle
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Adm12154 Kaposis s
Aad08215 Human gen
Adm12155 Herpesvir
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10.5	10.5	10.5	10.5	•	10.5	•	10.5	10.5	•	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5
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Adl67153 Plasmid p	Adl67175 Plasmid p	Aat40348 Plasmid p			Adl67148 Plasmid p		Adl67152 Plasmid p	Aa::22248 Nucleotid	Adl67154 Plasmid p	Abs66453 Plasmid p	Abs71027 pcsp-xa-F	Aav21683 Vector pl	Adp64415 Vector pC	Adm10659 Expressio	Aaz23778 Vector pS	Aax.90923 Anti-sens	Aai64275 Epstein-B	Aaa75454 Nucleotid	Adk65580 Human her	Aaf82902 EBV tethe	Aaa50254 Epstein B	Aax90924 Epstein B	Aav55831 Nucleotid	Aav55830 FLGA inse

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RESULT 1
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20-APR-2001
                                                                WPI; 2001-080711/09.
P-PSDB; AAB48317.
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                                                                                                                                                                                                                                            T. pallidum (CDC-2) acidic repeat protein (arp) encoding DNA
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                                                                                     Liu H,
                                                                                                                  14-JUN-1999;
                                                                                                                                14-JUN-2000; 2000WO-US016425.
                                                                                                                                                21-DEC-2000.
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Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological

The invention relates to a method of detecting presence of Treponema

Claim 19; Fig 7; 73pp; English.

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JOURNAL
MEDLINE
Query Match

Best Local Similarity

Matches 155; Conserv
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JOURNAL
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                                                                                                                                                                                                                                                                                                             alicat@sanger.ac.uk
see http://www.ebi.ac.uk/parasites/leish.html
betalls of Leishmania sequencing at the Sanger Centre are
at http://www.sanger.ac.uk/Projects/L major/
The cLHYG t3Hyg primer sequence can be obtained from acc.
                                                                                                                                                                                                                                                                                                                                                                                                                          2 (bases 1 to 67)
Taylor,R.G., Huckle, E.E.J., Ivens, A.C., Rajandream, M.A. and
Taylor,R.G.,
Barrell, B.G.
Direct Submission
Submitted (14-MAR-2000) Leishmania major Friedlin genome sequencing
project Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 365)
Ivens,A.C., Lewis,S.M., Bagherzadeh,A., Zhang,L., Chan,H.M. Smith,D.F.
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Leishmania major
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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nilarity 58.7%;
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AF342807 Treponema
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12.0	12.0	12.0	12.0	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.1	12.2	12.2	12.2	12.3	12.3	12.3	12.3	12.3	12.3
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		PEATURES
S OF AMERICA (US)	THE GOVERNMENT OF THE UNITED STATES	
	Patent: WO 0077486-A 3 21-DEC-2000;	JOURNAL
ting Treponema pallidum	Compositions and methods for detecting	TITLE
	Liu, H., Steiner, B. and Rhodes, B.	AUTHORS
		REFERENCE
Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.	Bacteria; Spirochaetes; Spirochaet	ONGREEN
	Treponema pallidum	SOURCE
•		KEYWORDS
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	·	ACCESSION
UNA LINEAR PAI 19-JAN-2001	Sequence 3 from Patent W00077486	DEFINITION
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                933.4
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76.8 1189
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AF411124 Treponema
AF411124 Treponema
AX068032 Sequence
AF015824 Treponema
AE001120 Treponema
AE01126 Treponema
AX068034 Sequence
AF342806 Treponema
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AC0125509 Mus muscu
AC0136718 Mus muscu
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12.7	12.7	12.7	12.9	13.3	13.7	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8
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Abk98592 Vector pE	Acd13882 L. lactis	Abk98631 Vector pE	Adq97246 Mouse can	Ach89664 Human gen	Aav55830 FLGA inse	Aax90924 Epstein B	Adn12161 Epstein-B	Adl71910 Expressio	Aaa59553 DNA clone	Ado07394 Modified	Ado07395 Modified			Adl67147 Plasmid p			Adl67153 Plasmid p	Adl67175 Plasmid p	Aat40348 Plasmid p		Ang51731 Plasmid p	Adl67148 Plasmid p	Adl67150 Plasmid p	ndl67152 Plasmid p

ALIGNMENTS

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RESULT 1

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AAC84649;

XX

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AAC84649;

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DT

11-SEP-2003 (revised)
DT

20-APR-2001 (first entry)

XX

Treponema pallidum; acidic repeat
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Treponema pallidum; ssp. endemicum
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Treponema pallidum; ssp. endemicum
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Treponema pallidum; ssp. endemicum
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21-DEC-2000; 2000WO-US016425.

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Detecting Treponema pallidum in bl

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Formation of the bacterium and an ar

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Sample.

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Claim 19; Fig 9; 73pp; English.

XX

CT

The invention relates to a method
                                                                                                             Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T. pallidum (Bosnia) acidic repeat protein (arp) encoding DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treponema pallidum; ssp. endemicum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treponema pallidum; acidic repeat protein; arp; immunogenic; syphilis; yaws; bejel; ds.
                                                                                                                                                                                                                                                                                                                                                                          (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
1. .939
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The invention relates to a method of detecting presence of Treponema

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AG433456 Mus muscu
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	r BAI UKES							COMPLETE	COMMENT			UCURNAL	TITLE	AUTHORS	REFERENCE	JOURNAL	TITLE	REFERENCE		ORGANISM	SOURCE	KEYWORDS	ACCESSION	DEFINITION	rocus	RESULT 1 AG561854/c	
/oz /mc /dt /cl /cl /cl	1776	R.Site 1 : ECORI R.Site 2 : ECORI	LIBRARY Vector : pBACe3.6	Sequencing : T7	e-mail: abe@rtc.riken.jp	Koyadai, Tsukuba, 305-0074 Japan	Tsukuba institute, bio kesource center, The Institute of Physical and Chemical Research (RIKEN) 3-1-1	library availability, please contact Kuniya Abe (abe@rtc.riken.jp).	Tel:81-45-503-9111, Fax:81-45-503-9170)	(E-mail:hattori@gsc.riken.jp, URL:http://hgp.gsc.riken.go.jp/,	1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan	and Chemical Research (RIKEN), Genomic Sciences Center (GSC);	Direct Submission	Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y.	2 (bases 1 to 776)	Unpublished	BAC end Seguences of Library MSMq01	I Colored to Manager to the Colored	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleoscomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.	Mus musculus molossinus	Mus musculus molossinus		AG561854 1 GT.48332552	Mus musculus molossinus DNA, clone:MSMg01-481H05.T7, genomic survey	AG561854 776 bp DNA linear GSS 05-JUN-2004		

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	CDS	gene		source	FEATURES	JOURNAL	TITLE	REFERENCE	JOURNAL		TITLE	AUTHORS	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	RESULT 1 AF411126	
/gene="arp" /codon_start=1 /transT_table=11 /transT_eacidic_repeat_protein" /product="acidic_repeat_protein" /protein_id="AAL07373.1" /db_xref="GI:15778315" /db_xref="GI:15778315" /translation="MFVRSDMFPKNTAVBISNLEKNAKAQAVVIGHAGIPGLLVSLAP /translation="MFVRSDMFPKNTAVBISNLEKNAKAQAVVIGHAGIPGLLVSLAP AAAAQLGIGVQAVRVRVRTLGTVRGGSGTSVGDUSLASLPSRVPAAPAQROPLSSPP AAAAQLGIGVQAVPRVRVRTLGTVRGGBREVEDVPKVVBPASEREGGEREVEDVPKV VEPASEREGGEREVEDVPKVVEPASEREGGEREVEDVPKVVEPASEREGGEREVEDVP	/gene="arp" 1. 1182	/sub_species="pertenue" /db_xref="taxon:168" 1.	<pre>/organism="Treponema pallidum subsp. pertenue" /mol_type="genomic DNA" /strain="CDC1"</pre>			-2001) National Center for	Direct Submission	This is a stained in the stained of	Unpublished	of Treponema pallidum	Molecular characterization of the acidic repeat protein gene (arp)	Liu, H., Steiner, B.M. and Rodes, B.	Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema. 1 (bases 1 to 1182)	Treponema pallidum subsp. pertenue	Treponema pallidum subsp. pertenue (yaws treponeme)		AF411126.1 GI:15778314	protern (arp) gene, comprete cas. AP411126	Treponema pallidum subsp. pertenue strain CDC1 acidic repeat			

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1101 684 974	1656 783 548 1100	992 1110 1448 870	1945 1560 1101 1231	776 709 697	Length
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				AG561854 Mus muscu AG351718 Mus muscu BY752869 BY752869 AZ337339 1M0068B03	Description
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FEATURES

Vector R.Site R.Site

: pBACe3.6 : EcoRI : EcoRI.

location/Qualifiers

Sequencing : T7 PRIMERS

JIBRARY

e-mail: abe@rtc.riken.jp

Bource

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tissue_type="mixture of kidney and spleen"/clone_lib="MSMg01 Mouse Male BAC Library"

'sex="male"

COMMENT

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AG424774 Mus muscu CB905422 tric074xf	-	AZ895651 RPCI-24-2	Dro	AG382169 Mus muscu	AG435295 Mus muscu	AG382042 Mus muscu	BP256465 HVSMEf001	CI,492176 SAIL 564	CL073978 CH216-131	CG748176 P042-1-A0	CG753585 P048-3-H1	AL,196350 Tetraodon	BX:998216 Reverse s	CL079800 CH216-156	AG448606 Mus muscu	CR.080757 Forward s	AZ383375 1M0141E03	CL058545 CH216-88E	CF877385 tric075xn

REFERENCE AUTHORS TITLE JOURNAL REFERENCE AUTHORS SOURCE ORGANISM ACCESSION VERSION KEYWORDS RESULT 1 AG561854/c LOCUS DEFINITION JOURNAL E 2 (bases 1 to 776) E 3 Hattori,M., Toyoda,A., Noguchi,H., Kojima,T. and Sakaki,Y. Birect Submission L Submitted (17-NOV-2003) Masahira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehiro-chou,Tsurumi-ku, Yokohama, Kanaguwa 230-0045, Japan (B-mail.hattori@gsc.riken.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-45-503-911, Fax:81-45-503-9170) Clones are derived from the mouse BAC library MSMg01. For BAC library availability, please contact Kuniya Abe (abe@rtc.riken.jp). The Institute of Physical and Chemical Research (RIKEN) 3-1-1 Koyadai, Tsukuba, 305-0074 Japan phone: 81-298-36-9189, fax: 81-298-36-9199 776 Mus musculus molossinus DNA, sequence. AG561884 Mus musculus molossinus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus Hattori,M., Toyoda,A., Noguchi,H., Kojima,T. and Sakaki,Y. BAC end Sequences of Library MSMg01 Unpublished Mus musculus molossinus AG561854.1 GI:48322552 bp DNA linear GSS 05-JUN-2004 clone:MSMg01-481H05.T7, genomic survey

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ACD13882	ABK98631	ADQ97246	AAX90924	AAK72613	ADN12161	ADL71910	AAA59553	ADO07394	AD007395	ADR12379	ADL67176	ADL67147	ADL67149	ADL67151	ADL67153	ADL67175	AAT40348	AAX15650	AAQ51731	ADL67148	ADL67150	ADL67152	AAZ22248	ADL67154
Acd13882 L. lactis	Abk98631 Vector pB	Adq97246 Mouse can	Aax90924 Epstein B	Aak72613 Human imm	Adn12161 Epstein-B	Adl71910 Expressio	Ana59553 DNA clone	Ado07394 Modified	Ado07395 Modified	Adr12379 Vector pC	Adl67176 Plasmid p	Adl67147 Plasmid p	Adl67149 Plasmid p	Adl67151 Plasmid p	Adl67153 Plasmid p		Aat40348 Plasmid p	Amx15650 Nucleotid	Aaq51731 Plasmid p	Adl67148 Plasmid p	Adl67150 Plasmid p		Auz22248 Nucleotid	Adl67154 Plasmid p

ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polynucleotide sequence from the genome of Treponema pallidum.
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AAX20500-21243 represent polynucleotide sequences from the genome of Treponema pallidum. The sequences can be used for detection, diagnosis, characterisation, prevention and therapy for T. pallidum infections, particularly syphilis. They can also be used for detecting diseases related to Borrelia infections in animals, and for the production of

Sequence 9410 BP; 1934 A; 2470 C; 2830 G; 2153 T;

0 U;

23 Other;

biosynthetic products such as enzymes

New isolated Treponema pallidum nucleic acids - used to develop products for the detection, diagnosis, characterisation, prevention and therapy of T. pallidum infections, particularly syphilis.

WPI; 1999-081273/07.

Fraser

Claim 1; Page 693-698; 1150pp; English.

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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES	

	AC122020	10	165592	11.5	148	19
AC127348 Mus	AC127348	10	229190	11.5	148.6	. 18
AC149611 Mus	AC149611	10	217353	11.5	148.6	c 17
AC116727 Mus muscu	AC116727	N	163945	11.5	148.6	c 16
AC136729 Mus muscu	AC136729	10	216602	11.6	149.6	c 15
AC073495 Mus muscu	AC073495	N	318930	11.7	150.2	C 14
AC096018 Rattus no	AC096018	Ŋ	260625	12.0	154	c 13
AC135961 Mus muscu	AC135961	10	197411	12.0	154.4	12
AC122509 Mus muscu	AC122509	10	178757	12.7	163	c 11
AC111089 Mus muscu	AC111089	N	211580	13.2	169.4	10
AX068034 Sequence	AX068034	_.	699	48.7	626.4	φ
AF342806 Treponema	AF342806	۳	1047	61.9	797	8
AF015824 Treponema	AF015824	۲	2946	69.7	896.4	7
AX068032 Sequence	AX068032	σ	2945	69.7	896.4	σ
AF411124 Treponema	AF411124	۳	1647	70.6	909	Ų
AX068036 Sequence	AX068036	6	939	72.5	933.4	4
AF411126 Treponema	AF411126	1	1182	82.9	1067	ω
AE001220 Treponema	AE001220	<u>مــر</u>	14268	87.4	1125	N
AF342807 Treponema	AF342807	۳	1287	100.0	1287	_
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11.0	11.1	11.1	11.1	11.1	11.1	11.1	11.1	11.2	11.2	11.2	11.2	11.2	11.2	11.2	11.3	11.3	11.3	11.3	11.3	11.3	11.3	11.5	11.5
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	source	FEATURES	COMMENT		JOURNAL	AUTHORS	REFERENCE		JOURNAL	TITLE	AUTHORS	REFERENCE	JOURNAL		TITLE	AUTHORS	REFERENCE	JOURNAL		TITLE	AUTHORS	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	RESULT 1 AF342807
/organism="Treponema pallidum subsp. endemicum" /mol type="genomic DNA" /strain="Bosnia" /sub_species="endemicum" /db_xref="taxon:53436"	11287		Sequence update by submitter On sen 11 2001 this semience version replaced of:12667500.	ISA	Submitted (13-SEP-2001) National Center for Infectious Diseases,	Liu, H., Steiner, B. and Rodes, B.	4 (bases 1 to 1287)	Frevencton,	Submitted (25-JAN-2001) National Center for Intectious Diseases,		Liu, H. and Steiner, B.	3 (bases 1 to 1287)	Unpublished	Treponema pallidum	Molecular characterization of the acidic repeat protein (arp) of	Liu, H., Steiner, B.M. and Rodes, B.	2 (bases 1 to 1287)		subspecies endemicum (Bosnia strain)		Liu, H. and Steiner, B.	Dattelia; opilotiaetes; opilotiaetaies; opilotiaetaieac, inclosiess. 1 (bases 1 to 1287).				AF342807.2 GI:15617206	AF342807	Treponema paliidum subsp. endemicum strain Bosnia acidic repeat	linear	

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ALIGNMENTS

REFERENCE AUTHORS TITLE JOURNAL REFERENCE SOURCE ORGANISM ACCESSION VERSION KEYWORDS RESULT 1 AG561854/c LOCUS COMMENT FEATURES DEFINITION JOURNAL **Bource** Vector R.Site R.Site Direct Submission Submitted (17-NOV-2003) Masshira Hattori, The Institute of Physical Submitted (17-NOV-2003) Masshira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (SSC); 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan 16-8-81 institution of the State of the Sta library availability, please contact Kuniya Abe Tsukuba Institude, Bio Resource Center, The Institute of Physical and Chemical Research Koyadai, Tsukuba, 305-0074 Japan phone: 81-298-36-9189, fax: 81-298-36-9199 e-mail: abe@rtc.riken.jp Mus musculus molossinus Mus musculus molossinus Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus Mus musculus molossinus Sequencing : T7 LIBRARY Bequence. AG561854 PRIMERS 2 (bases 1 to 776) Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y. Hattori,M., Toyoda,A., Noguchi,H., Kojima,T. and Sakaki,Y. BAC end Sequences of Library MSMg01 AG561854.1 AG561854 Jnpublished N 14 /organism="Mus musculus molossinus" mol type="genomic DNA" /sub_species="molossinus" /db xref="taxon.57486" /clone="MSMg01-481H05.T7" /sex="male" Location/Qualifiers tissue_type="mixture of kidney and spleen" /clone_lib="MSMg01 Mouse Male BAC Library" : pBACe3.6 : EcoRI GI:48322552 EcoRI DNA, 776 bp DNA linear (clone:MSMg01-481H05.T7, library NSMg01. For BAC (uniya Abe (abe@rtc.riken.jp) (RIKEN) 3-1-1 GSS 05-JUN-2004 , genomic survey

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(first entry)
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1. .939
                                                                                                                                                                                                                                                                  immunogenic; syphilis;
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14-JUN-1999; 14-JUN-2000; 2000WO-US016425. 21-DEC-2000. WO200077486-A2 (USSH) US DEPT HEALTH & HUMAN SERVICES. 99US-0138981P.

Liu H, Steiner B,

Rhodes B;

Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological sample. WPI; 2001-080711/09. P-PSDB; AAB48318.

Claim 19; Fig 9; 73pp; English.

The invention relates to a method of detecting presence of Treponema

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Result
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AF342807 Treponema
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ALIGNMENTS

	source	FEATURES	REMARK		JOURNAL	TITLE	AUTHORS		COURTAIN	TITLE	AUTHORS	REFERENCE	JOURNAL	ŧ	TITLE	AUTHORS	REFERENCE	JOURNAL	11116	AUTHORS	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AF342806	RESULT 1
<pre>/organism="Treponema pallidum subsp. pe::tenue" /mol_type="genomic DNA" /strain="CDC2" /sub_species="pertenue" /db_xref="taxon:168"</pre>	11047	Location/Qualifiers		nd Prevention, USA	ional Center for	Direct Submission	4 (bases 1 to 1047) Liu,H., Steiner,B. and Rodes.B.	Stop D13, Atlanta, GA 30333, USA	ional Center for		Liu, H. and Steiner, B.	3 (bases 1 to 1047)	Unpublished	Treponema nallidum	The social conservation of the second		2 (bases 1 to 1047)	Unoublished	avidance repeat protein (arp) gene sequence of Treponema pallidum		1 (bases 1 to 1047)	pallidum subsp. pertenue			AF342806.2 GI:15617204		a pallidum subsp. pertenue strain CDC2 acidi	AF342806 1047 bp DNA linear BCT 13-SED-2001		

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Result
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          BG273097
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AG561854 Mus muscu
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AZ383375 1M0141E03
CD328191 AGENCOURT
                                             BG273097 nai99f06.
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4 OGUBU35TH
4 PUGAG07TB
8 OG2BH42TH
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6.5	6.5	6.5	6.5	6.5	6.5	6.5	6.5	6.5	6.5	6.6	6.6	6.6	6.6	6.6	6.6	6.6	6.6	6.6	6.6	6.7
889	651	651	647	624	609	592	552	520	501	1344	583	516	494	986	430	1271	699	1147	427	1197
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MEDLINE PUBMED REFERENCE AUTHORS ACCESSION VERSION KEYWORDS SOURCE ORGANISM LOCUS DEFINITION RESULT 1 L2644X/c ORIGIN REFERENCE FEATURES COMMENT TITLE JOURNAL TITLE AUTHORS alicat@sanger.ac.uk
see http://www.ebi.ac.uk/parasites/leish.html
betails of Leishmania sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/projecte/L_major/
The cLHYG t3Hyg primer sequence can be obtained from and no Submitted (14-MAR-2000) Leishmania major Friedlin genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and Direct Submission 2 (bases 1 to 67)
Taylor,R.G., Huckle, E.E.J., Ivens, A.C., Rajandream, M.A. and
Barrell, B.G. A physical map of the Leishmania major Genome Res. 8 (2), 135-145 (1998) Ivens,A.C., Lewis,S.M., Bagherzadeh,A.,
Smith,D.F. Leishmania major Eukaryota; Euglenozoa; Kinetoplastida; Trypanosonatidae; AL160994.1 GI:7258621 98146435 Leishmania major (bases 1 to 365) CLHYG t3Hyg primer sequence can be obtained from /db_xref="taxon:5664" /chromosome="26" /clone="cosmid L2644" /organism="Leishmania major"
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/strain="Friedlin" . .365 location/Qualifiers Friedlin Zhang, L., Chan, H.M.

Query Match 8.6%; Best Local Similarity 58.7%; Matches 155; Conservative

Score 89.6; DB 9; Pred. No. 1.5e-13; 0; Mismatches 109;

Length 365;

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Gaps

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Minimum DB seq length: 0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1 852.2 81.4 66.2 693.4 66.3 3 4 626.4 59.8 82.6 7.9 6 82.6 7.0 7.0 8 82.6 7.0 7.0 8 82.6 7.0 7.0 8 82.6 7.0 7.0 8 82.6 7.0 8 82.6 7.0 8 82.0 7.0 8 82.0 8 82.0 7.0 8 82.0 8 82.0 8 82.0 8 82.0 8 82.0 8 82.0 8 8
2 693.4 66 4 626.4 59 5 82.6 7 7 82.6
3 642 61 5 82.6 57 6 82.6 77 8 82.6 77 9 82.6 77 11 82.6 77 11 82.6 77 11 82.6 77 11 82.6 77 11 82.6 77 11 77.4 77 11 77.8 87 11 77.8 87 12 77.8 87 13 77.8 87 14 77.8 87 15 77.8 87 17 77.8 87 17 77.8 87 18 77.8 87 19 77.
4 626.4 59 6 82.6 7 7 82.6 7 8 82.6 7 9 82.6 7 11 82.6 7 11 82.6 7 11 82.6 7 11 77.4 7 11 77.4 7 11 75.8 7 11 75.8 7 75.8 7 75.8 7
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13 80 14 78 15 78 16 77 17 77 18 75 19 75
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16 77. 17 77. 18 75. 19 75. 20 7
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c 18 75.8 7 19 75.8 7 c 20 75 7
19 75.8 7 c 20 75 7
c 20 75 7

AAX20500-21243 represent polynucleotide sequences from the genome of Treponema pallidum. The sequences can be used for detection, diagnosis, characterisation, prevention at therapy for T. pallidum infections, particularly syphilis. They can also be used for detecting diseases related to Borrelia infections in animals, and for the production of biosynthetic products such as enzymes

Sequence 9410 BP; 1934 A; 2470 C; 2830 G; 2153 T; 0 U; 23 Other;

New isolated Treponema pallidum nucleic acids - used to develop products for the detection, diagnosis, characterisation, prevention and therapy of T. pallidum infections, particularly syphilis.

WPI; 1999-081273/07.

Claim 1; Page 693-698; 1150pp; English.

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73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4	73.4
7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0	7.0
10774	10615	10596	10596	10596	10561	10516	10477	10380	10330	10285	10285	9600	9482	8705	8705	5452	2580	2580	1926	1926	1926	1925	799	795
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ADL67153	ADL67175	AAT40348	AAX15650	AAQ51731	ADL67148	ADL67150	ADL67152	AAZ22248	ADL67154	ABS66453	ABS71027	AAV21683	ADP64415	ADM10659	AAZ23778	AAX90923	AAI64275	AAA75454	ADK65580	AAF82902	AAA50254	AAX90924	AAV55831	AAV55830
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RESULT 1
AAX20603
ID AAX2
XX
Treponema pallidum infection; syphilis; Borrelia infection; animal; enzyme production; ds.
                                                                                                           Fraser CM;
                                                                                                                                    24-JUN-1997;
                                                                                                                                                23-JUN-1998;
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                                                                                                                                                                                                                       Polynucleotide sequence from the genome of Treponema pallidum
                                                                                                                       (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                     (first entry)
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